```
logo

NTDB id 1051072 ACEWX4 RS12770 WP 168383358.1  MPMFKLLAHSHTLFKKLLPCQLCGSAAQQQYRVCADCLQQLPWLKASVCRQETEIQVACHYRYPVDRIIQQFKYEQQLHY   80
NTDB id 1071 ABD1 RS15375 WP 000472269.1       ..MFKFL.NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
NTDB id 1042 HON27 RS01930 WP 000472273.1      ..MFKFL.NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
consensus                                        !!!*! ****!!**!*!!*!!*******!**!**!**!!!!!!****!******!!!!!*!!**!!!!!!!!!!!*!!!

logo

NTDB id 1051072 ACEWX4 RS12770 WP 168383358.1  QSLLAGSLCELAYPRVQAIVPMPVAAERLAERGYHHTLLLAQVLARHLQLPIWQPVVRHHQHSQKGLSRLERLENLDQQF  160
NTDB id 1071 ABD1 RS15375 WP 000472269.1       QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
NTDB id 1042 HON27 RS01930 WP 000472273.1      QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
consensus                                      !*!!!**!**!**!*!!!!!!!!****!!*!!!****!!!!**!***!**!*!!!!*!***!!!!!!!!!!!!*!!**!!!

logo

NTDB id 1051072 ACEWX4 RS12770 WP 168383358.1  QLRPDTGRPYRRVLIVDDVVTTGSSIRALKQQLEQLGCHKVYAVCIASADG...  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1       VALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1042 HON27 RS01930 WP 000472273.1      VALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                      ****** !*!!!!!!!*!!!*!!!!!!*!!*!*!*!!!!****!*!*!*!*****
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X non conserved

X similar

X $\geq$ 50% conserved